GeneMark

Version 2.5p (09.08.06) *Copyright 1993 - M. Borodovsky, J. McIninch*

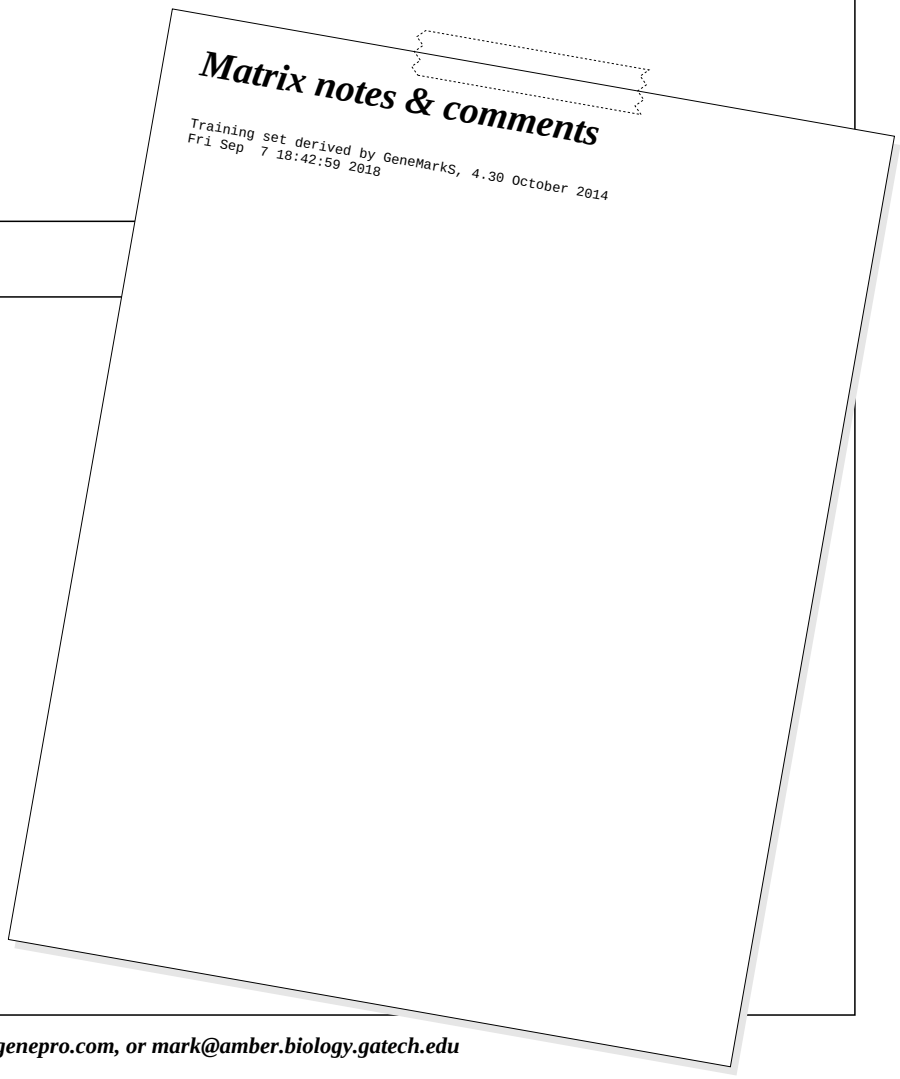
PROGRAM INFORMATION

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Sequence         :  Microbacterium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster
Analysis Date    :  3/28/24 at 10:22:50
Pages            :  21
Sequence Length  :  41770 bp
GC Content       :   63.44%

Window Length    :  96 bp
Window Step      :  12 bp
Threshold Value  :  0.500

PS-Version       :  1.2

GeneMark Options :  PostScript graph,
                    Mark ORFs / splice sites,
                    List ORFs,
                    List regions and/or splice sites,
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MATRIX INFORMATION

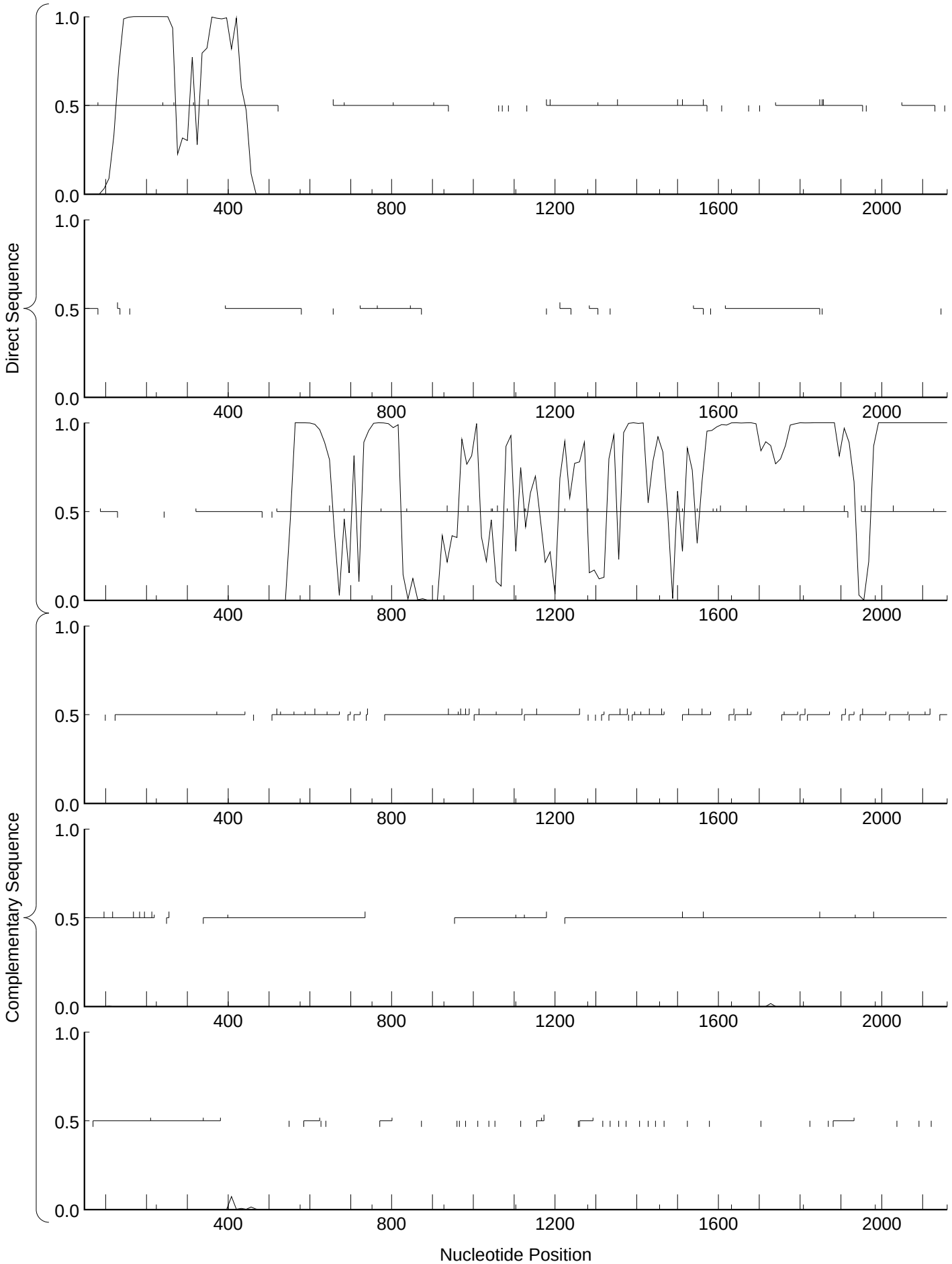
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Matrix     :  -
Author     :  -

Order      :  4
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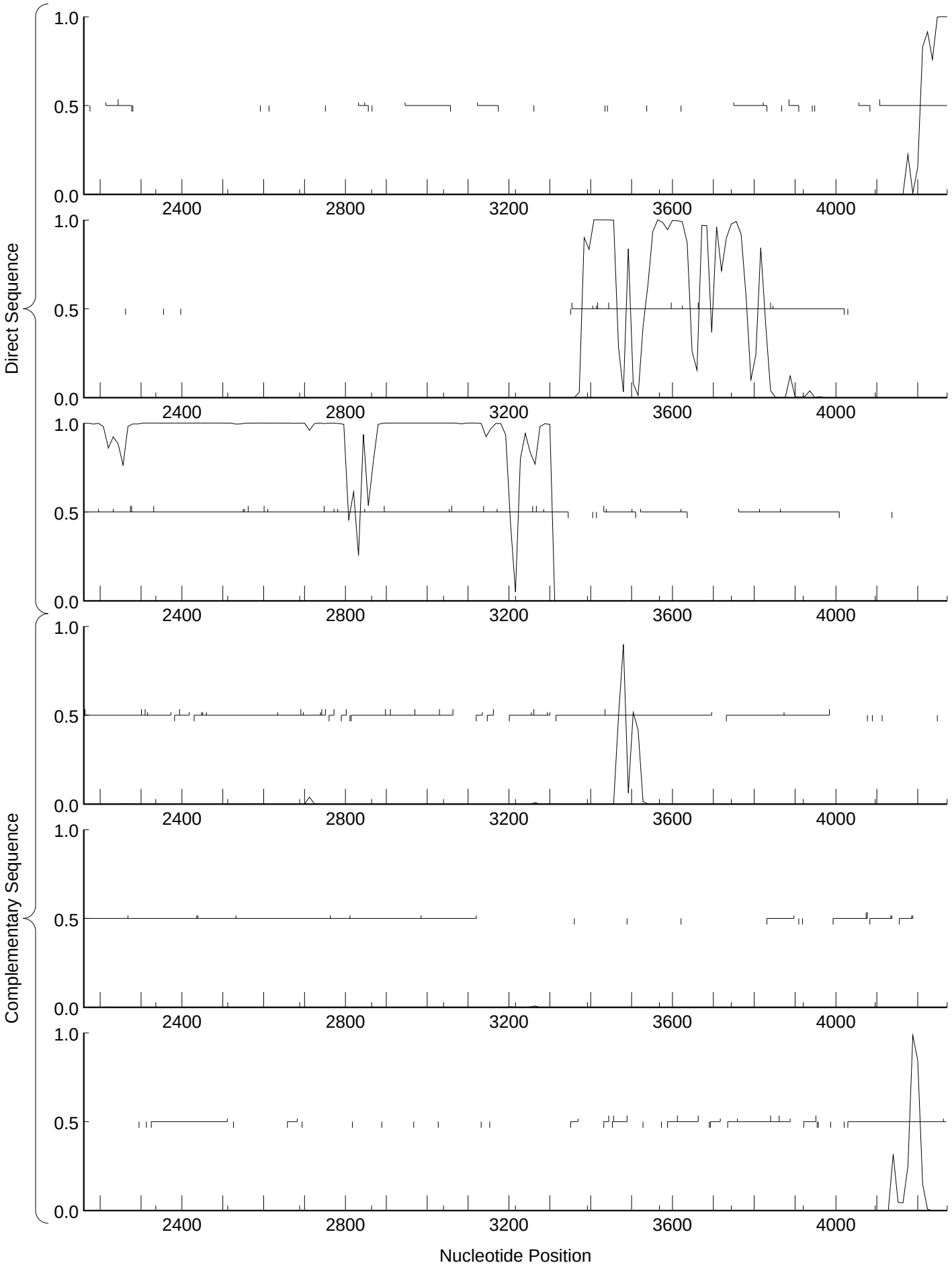

```
  Send questions / comments to:
      Dr. M. Borodovsky
      Georgia Institute of Technology
      School of Biology
      Atlanta, GA  30332-0230
```


For help or information, send electronic mail to: custserv@genepro.com, or mark@amber.biology.gatech.edu

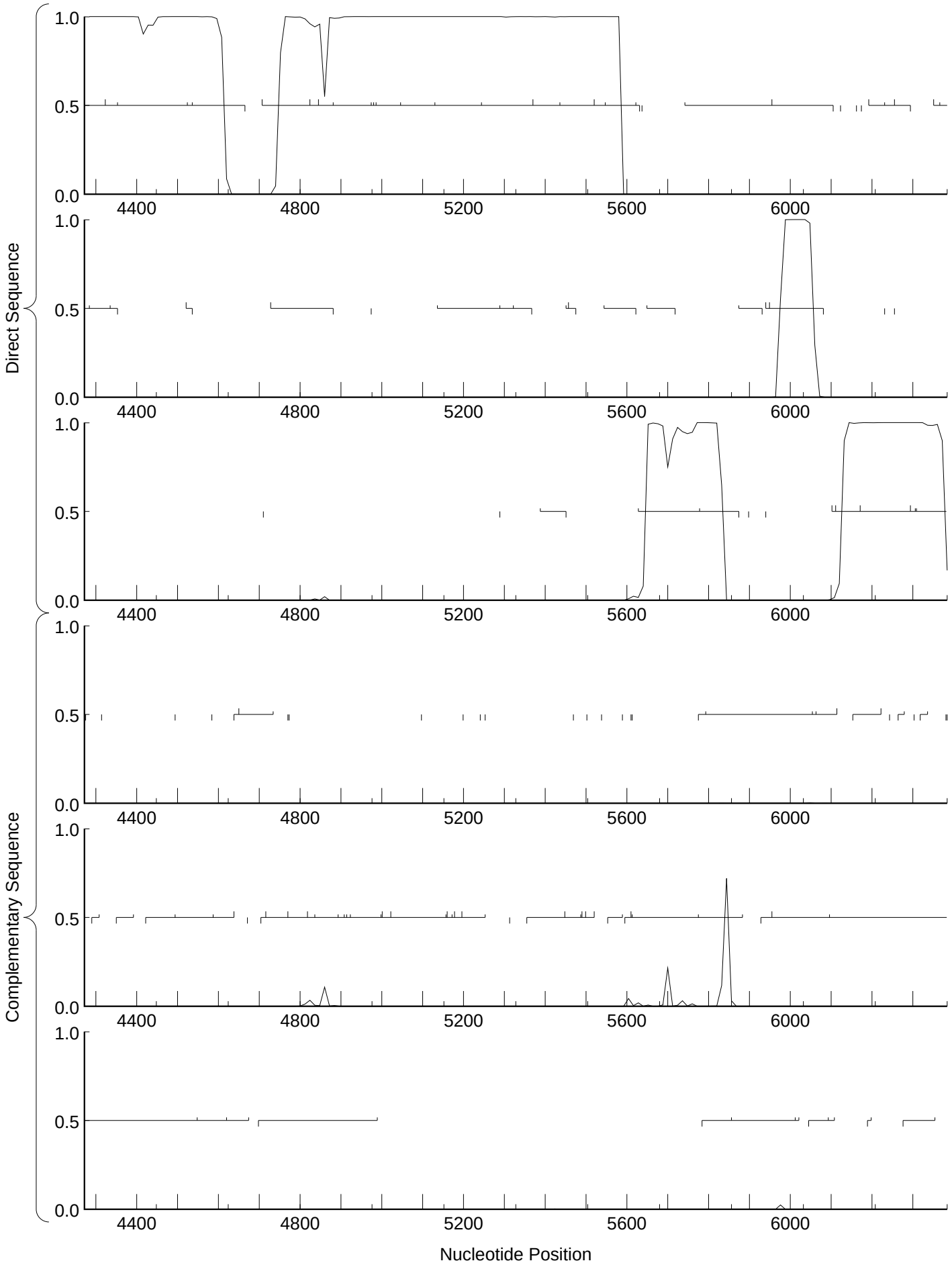
cterium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 2/21



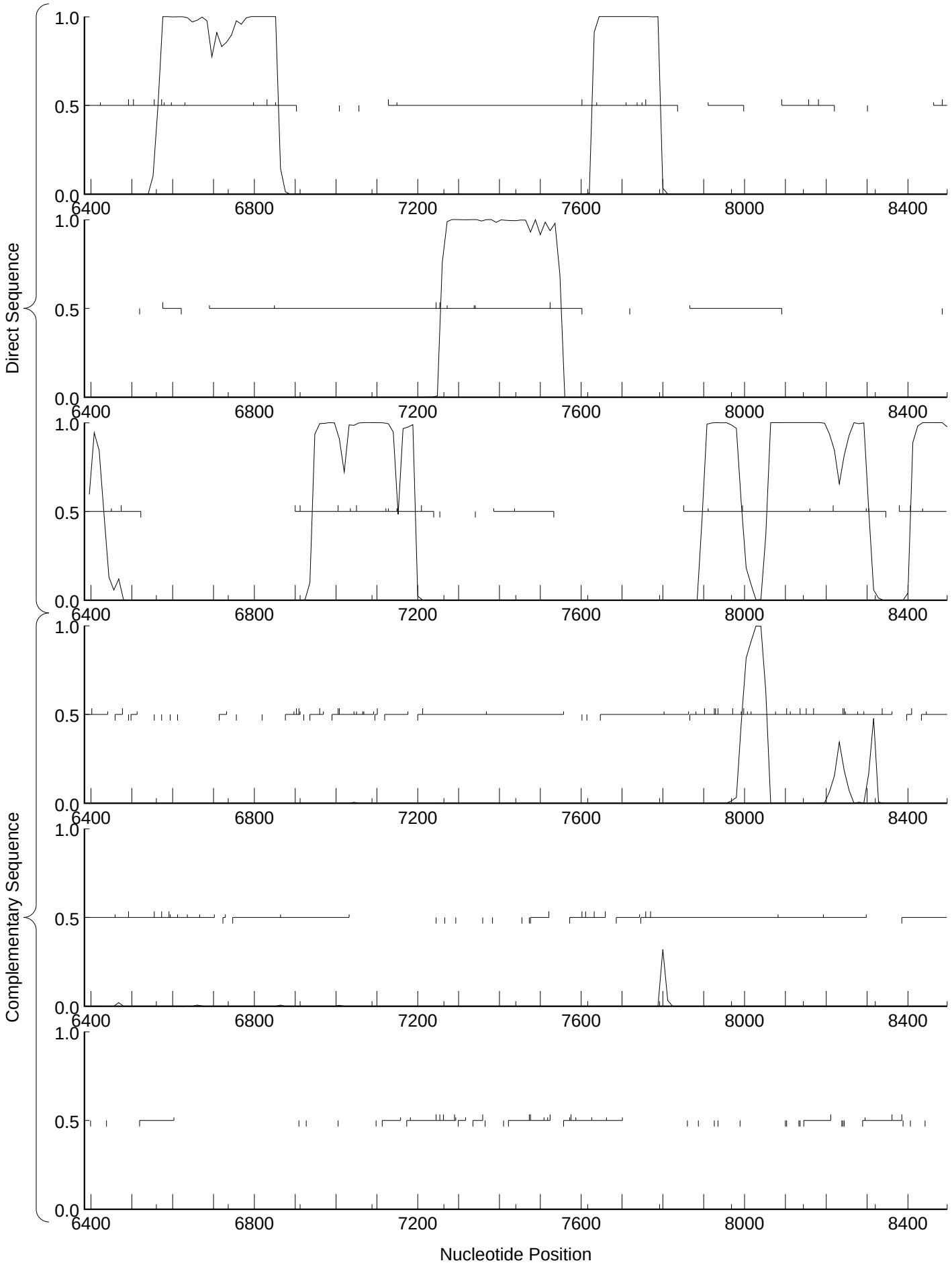
cterium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 3/21



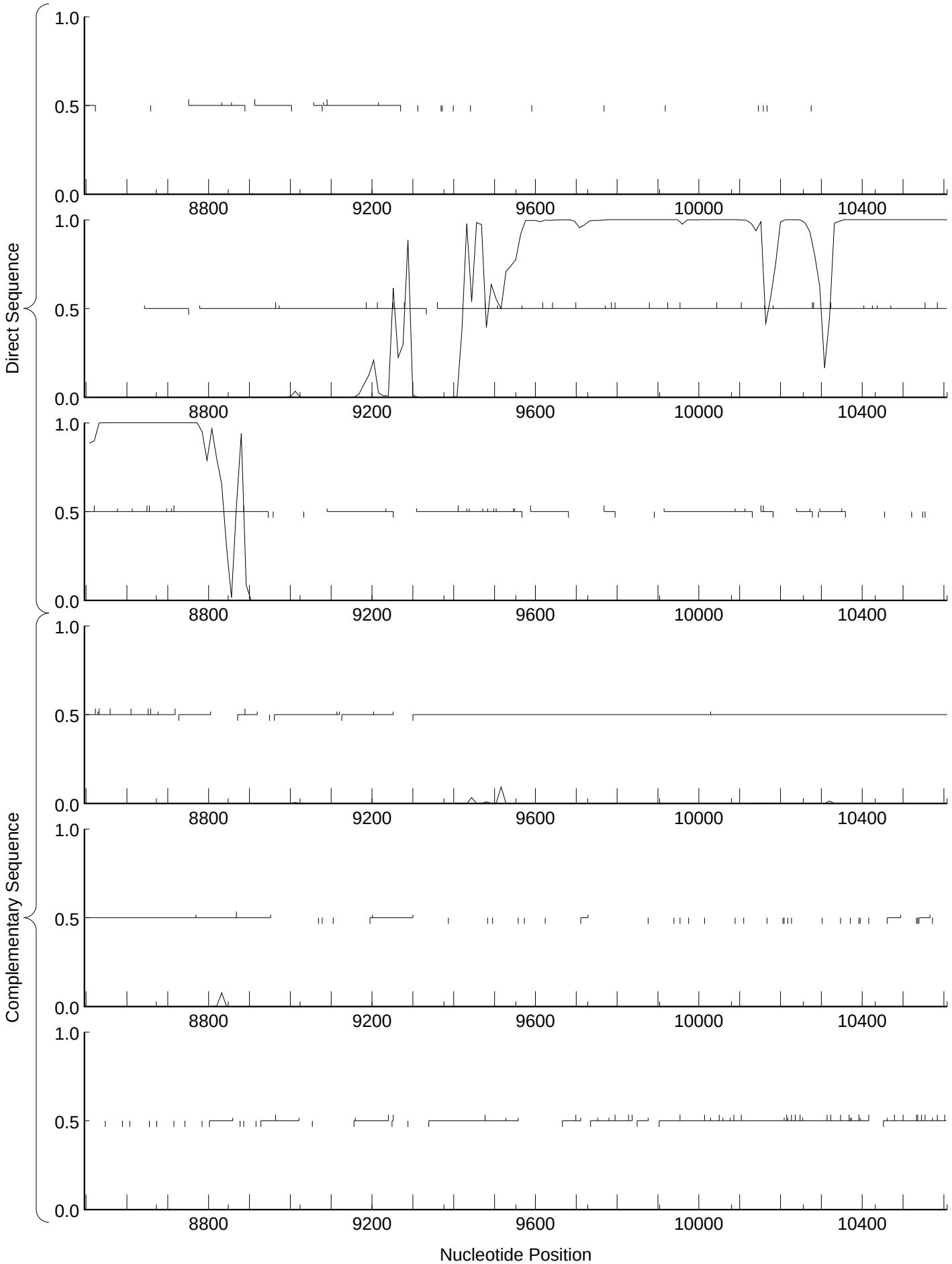
cterium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 4/21



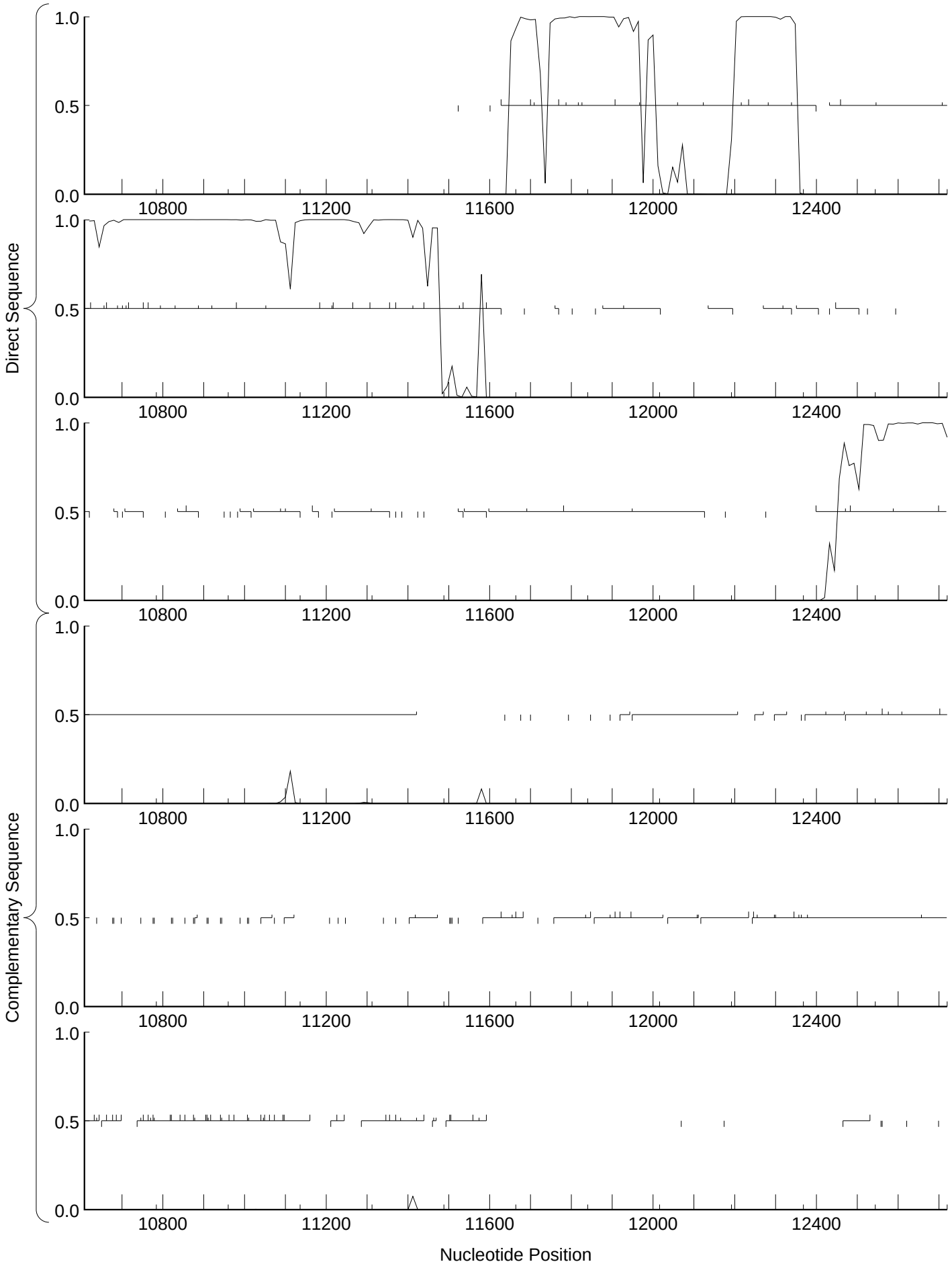
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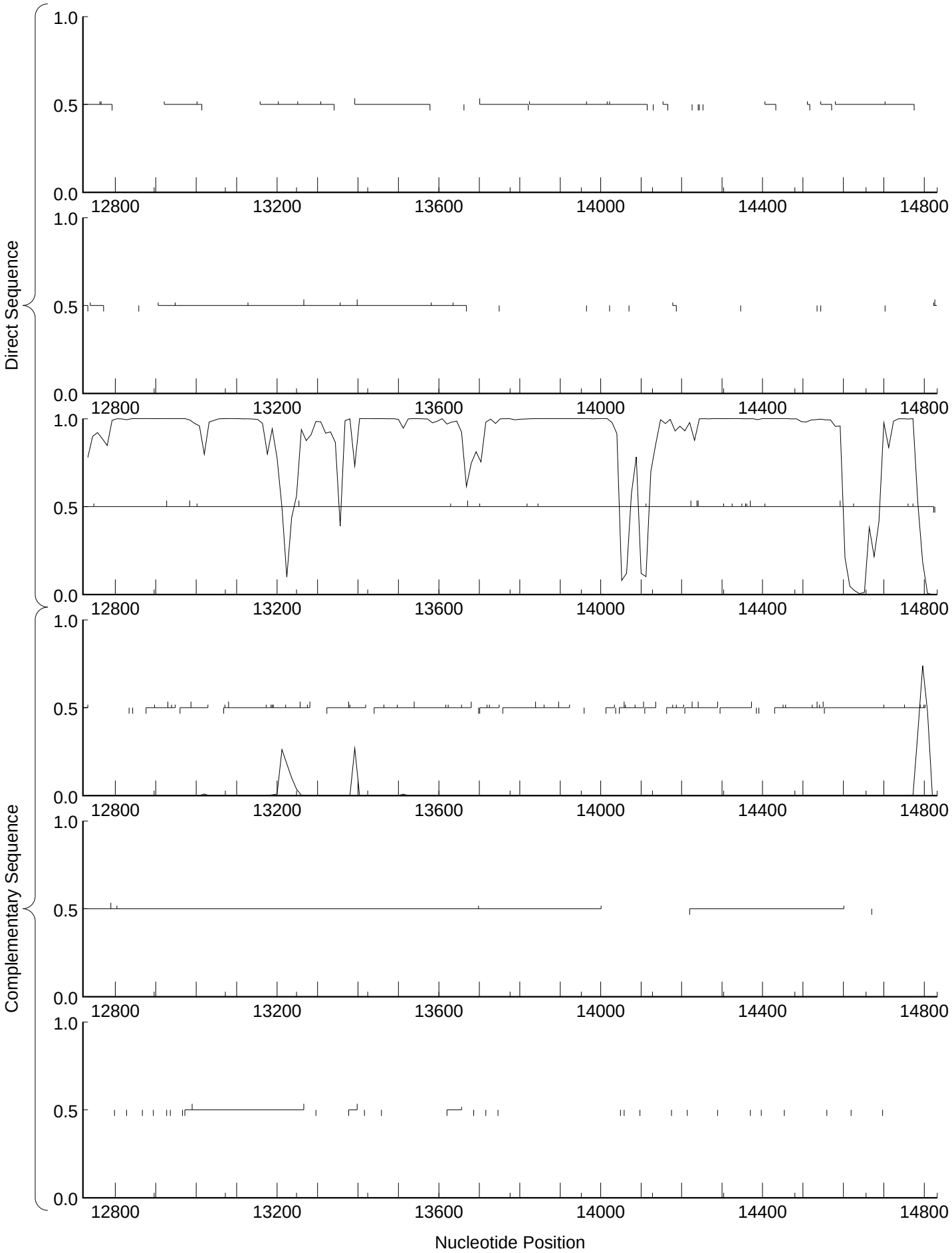
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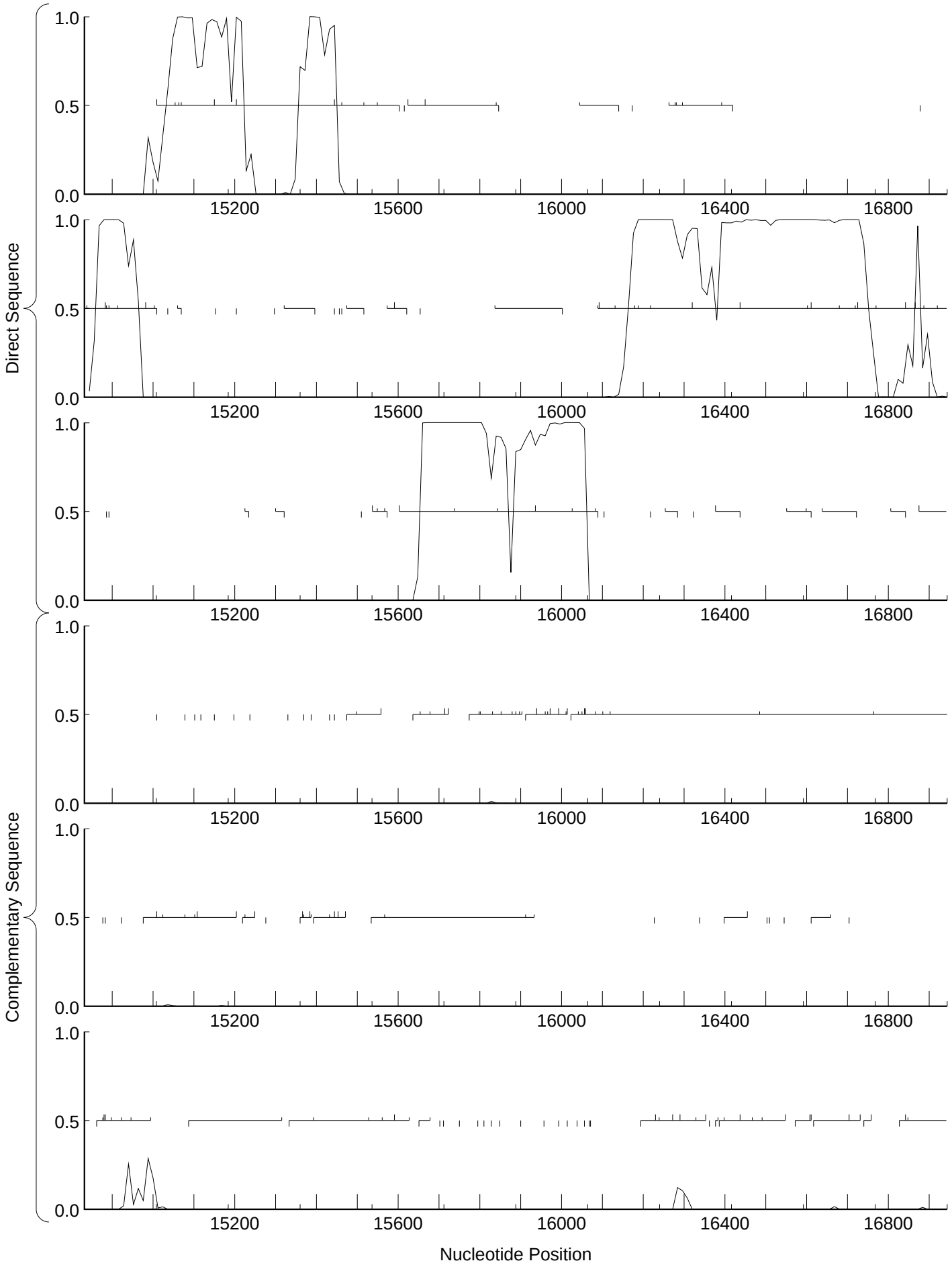
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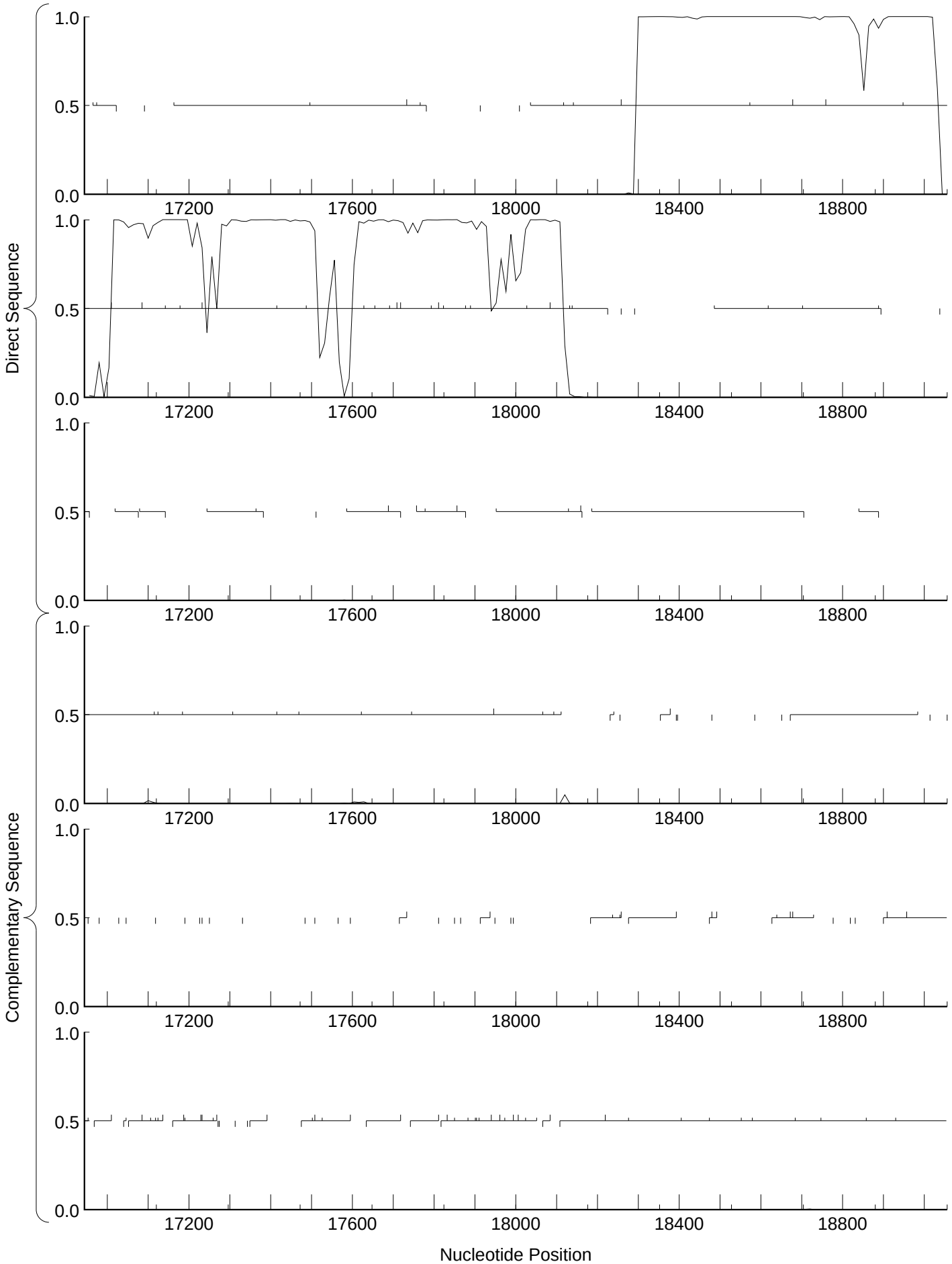
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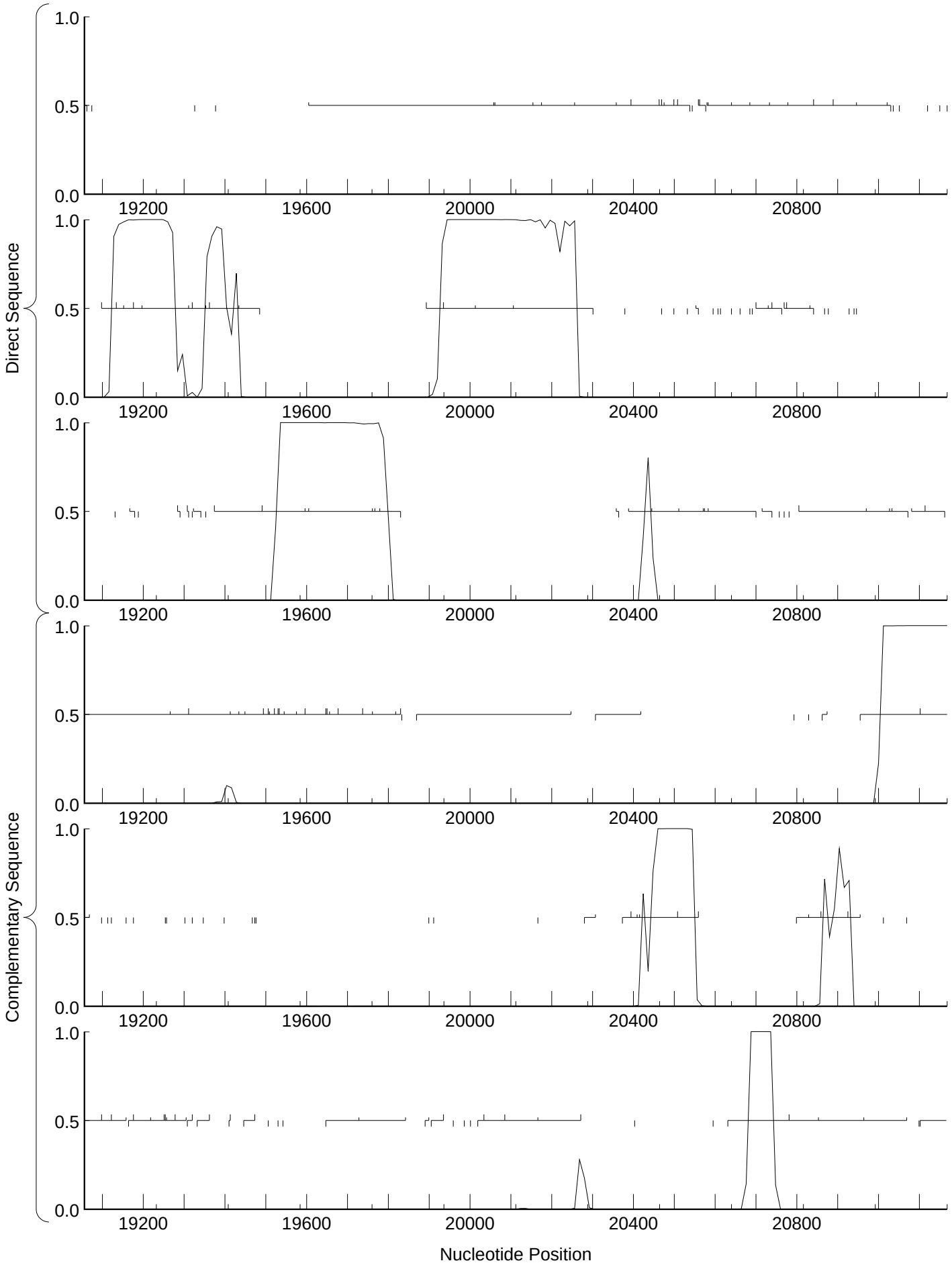
cterium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 9/21



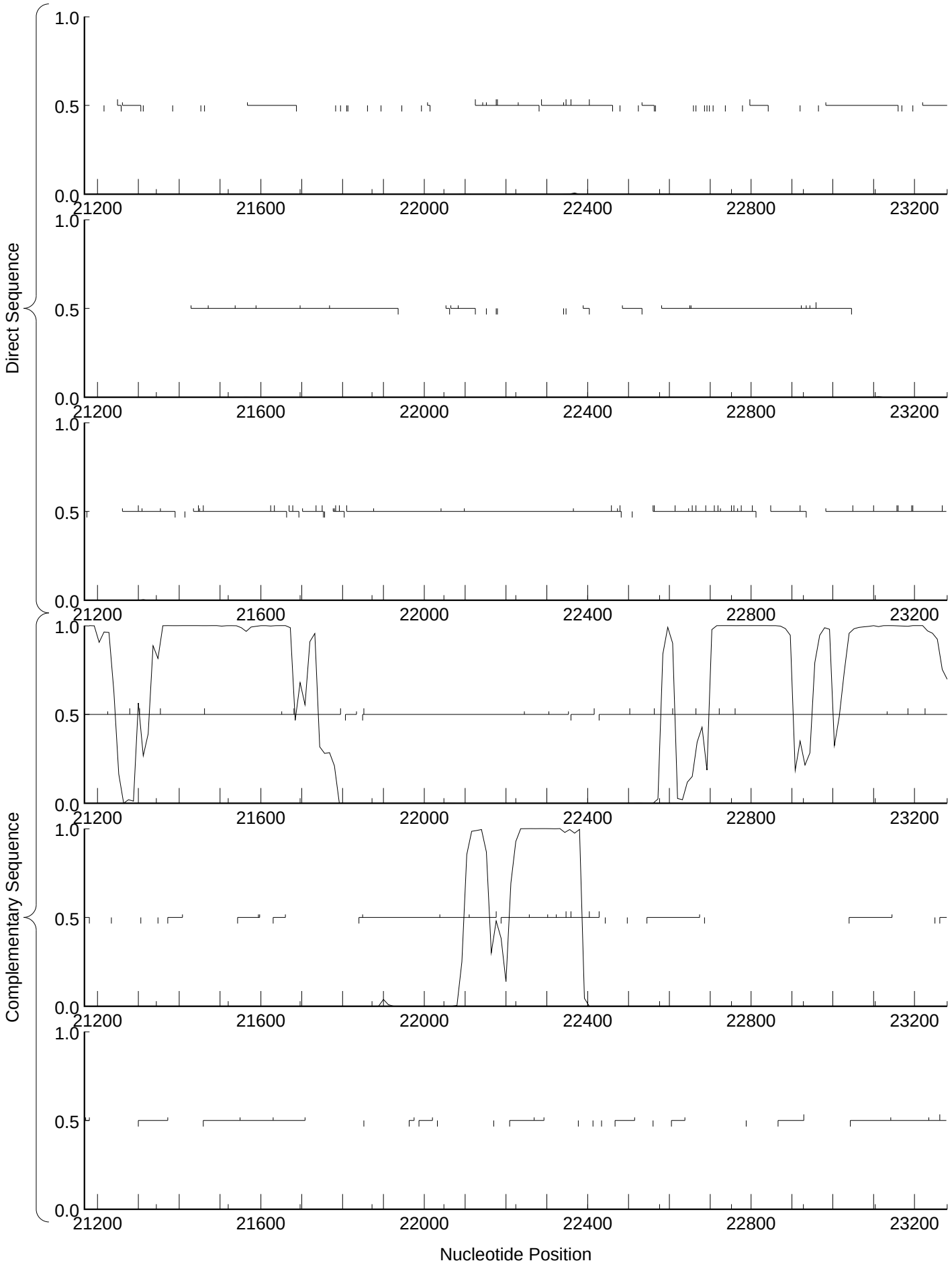
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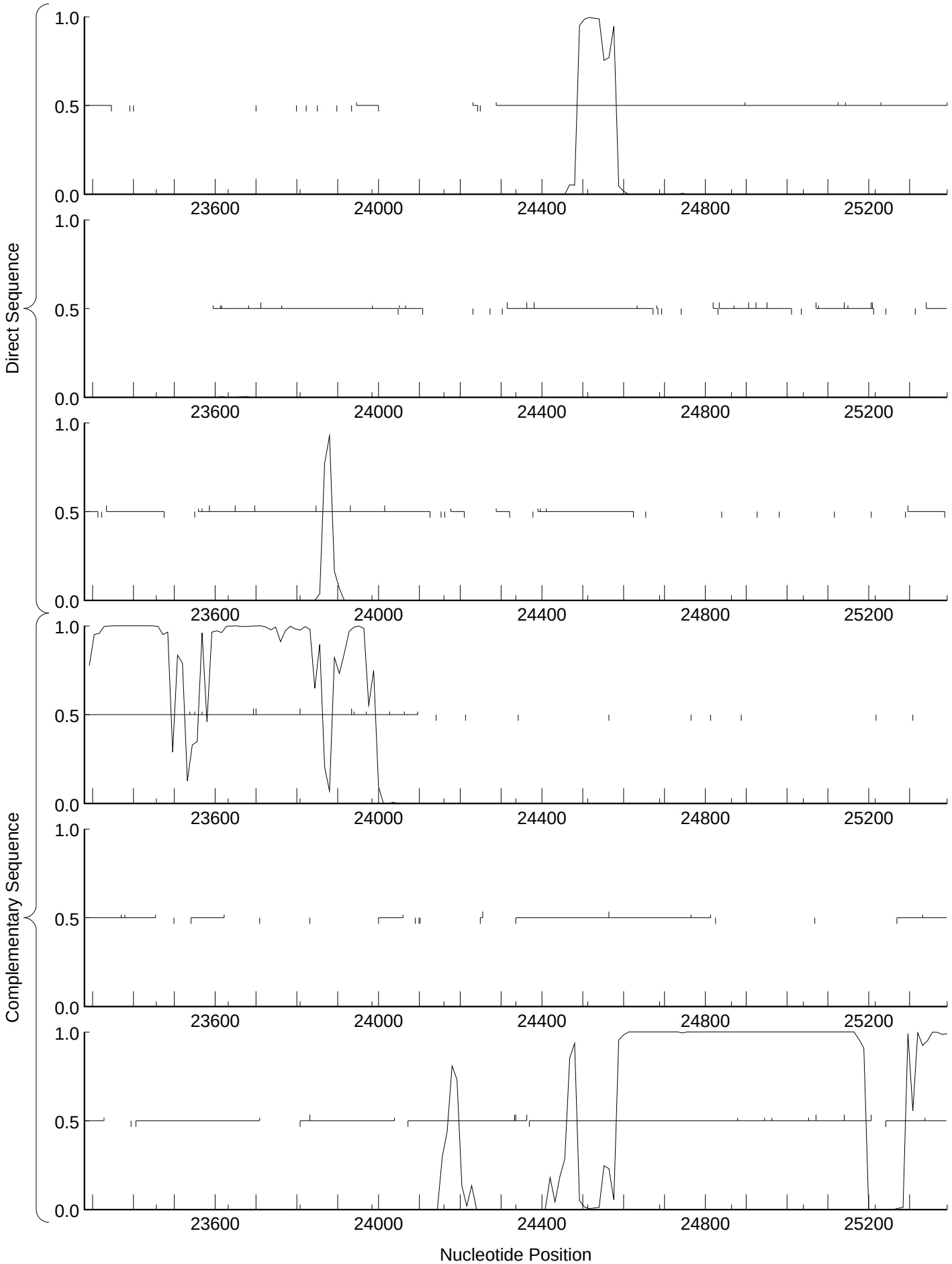
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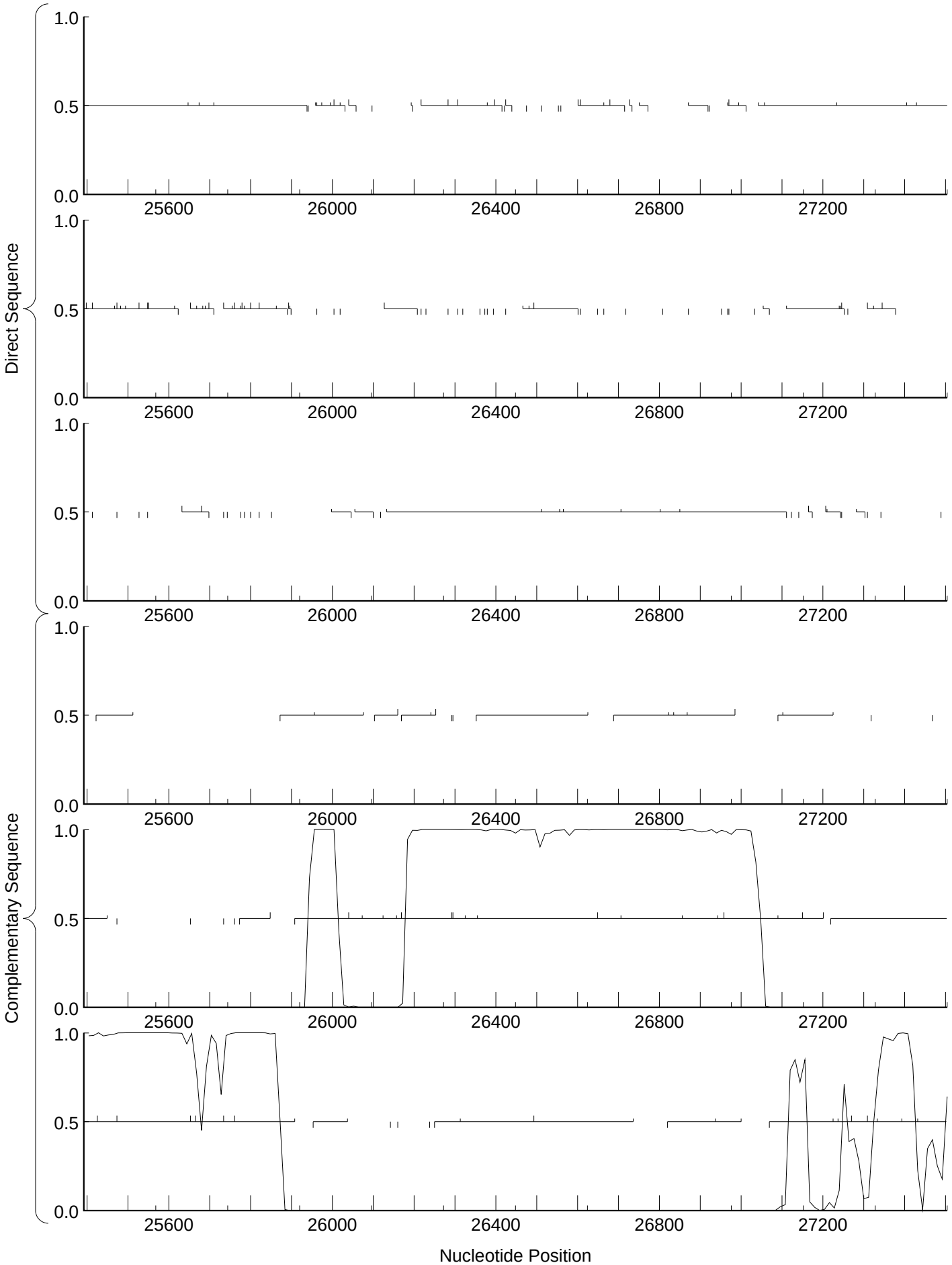
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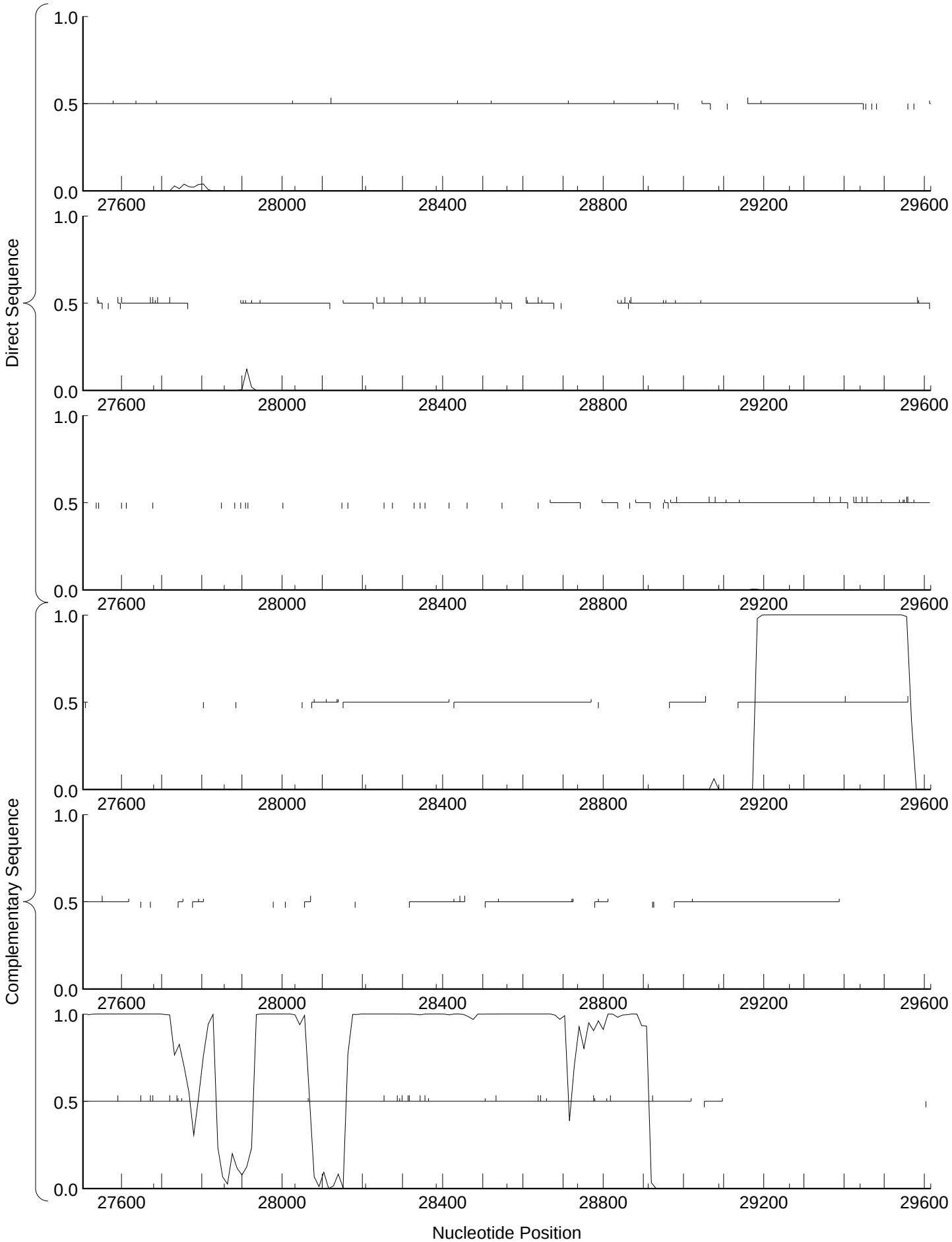
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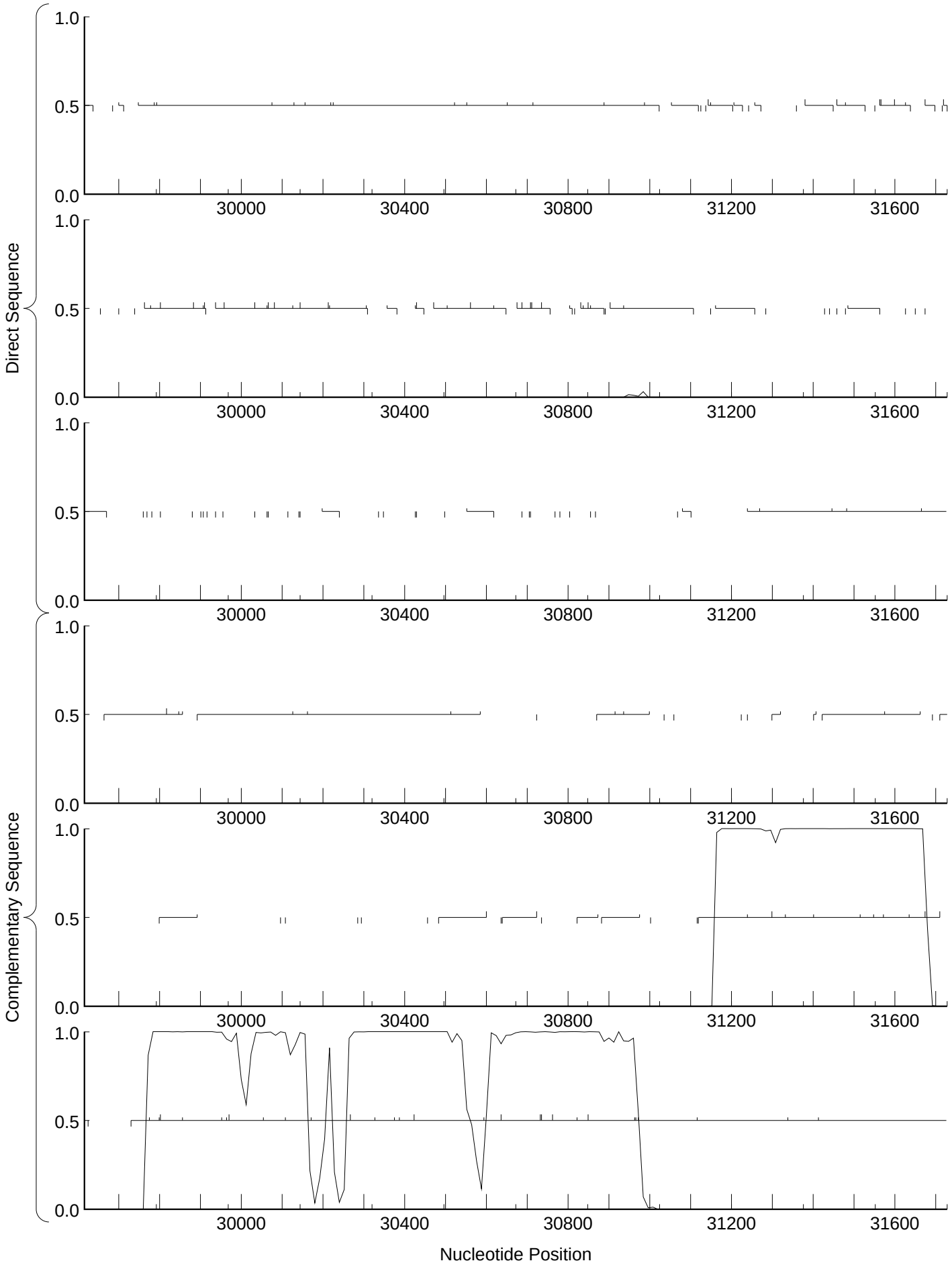


erium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 14/21

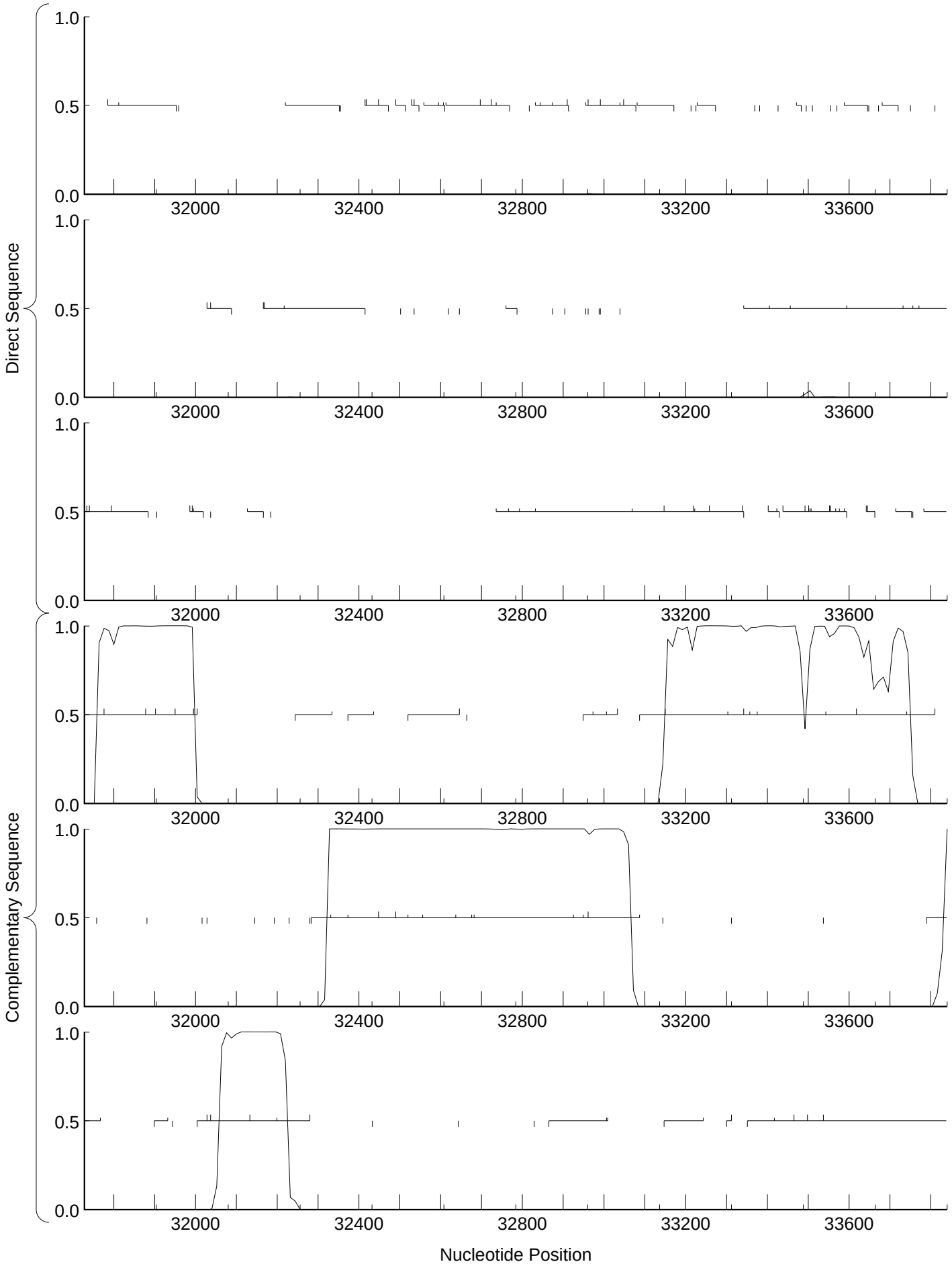


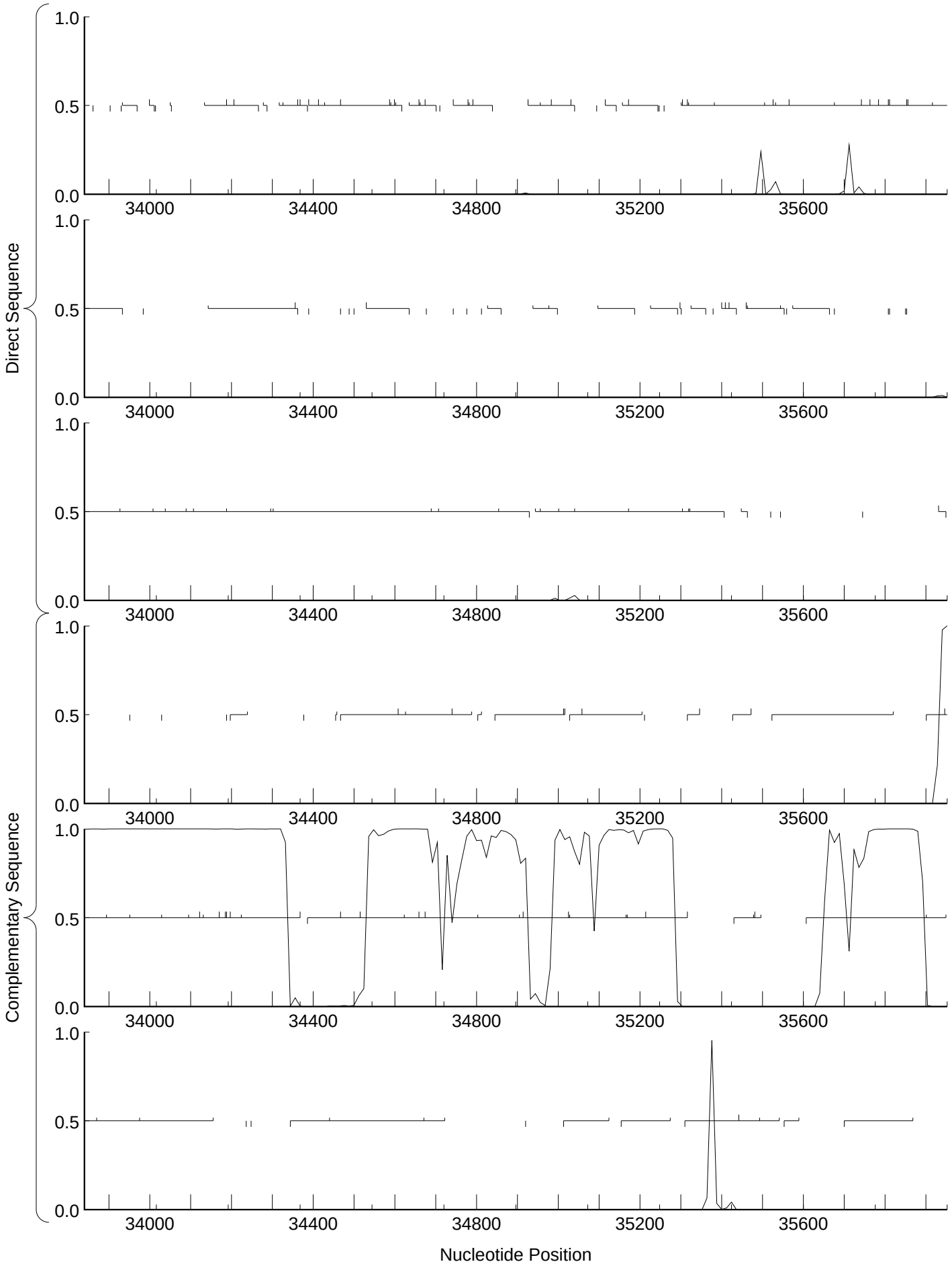
erium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 15/21



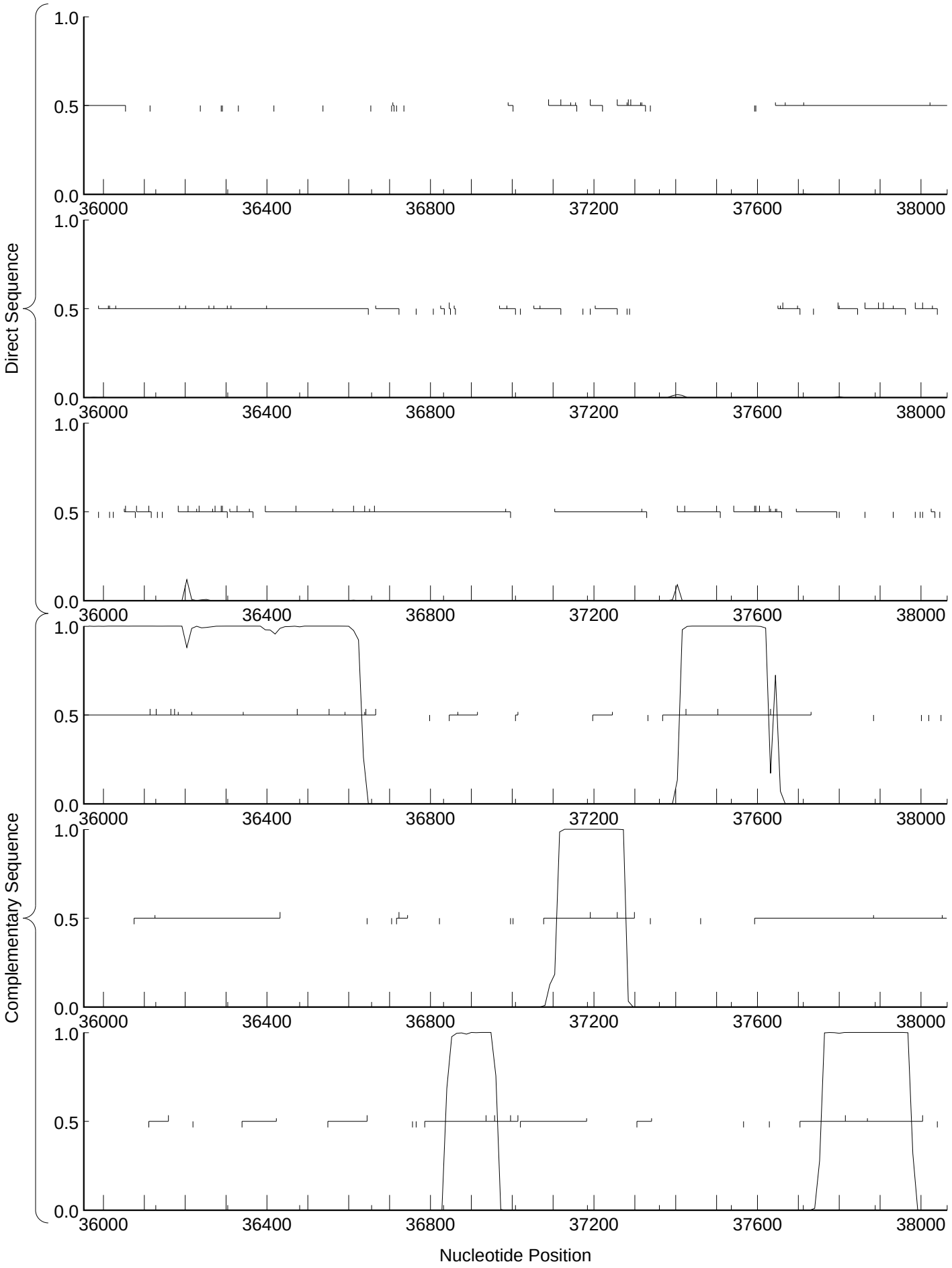

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Complementary Sequence
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0.5
0.0
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30400
30800
31200
31600
Nucleotide Position

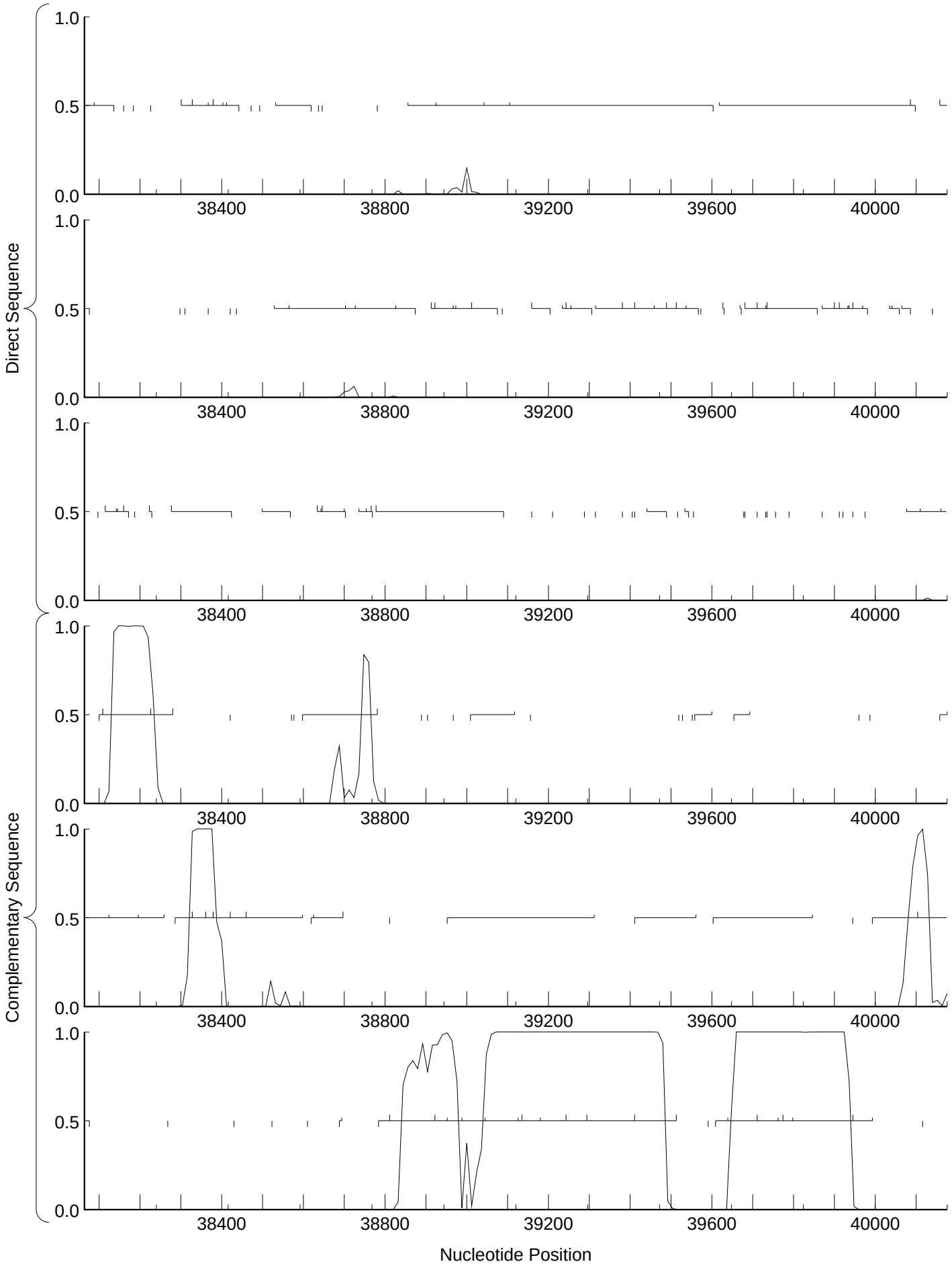
erium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 17/21




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
34000
34400
34800
35200
35600
Nucleotide Position

erium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 19/21




Direct Sequence
Complementary Sequence
1.0
0.5
0.0
38400
38800
39200
39600
40000
Nucleotide Position

erium phage Alyxandracam complete sequence, 41770 bp, circularly permuted, Cluster EA1, Order 4, Window 96, Step 12, 21/21

